

SEQUENCE LISTING

<110> INCYTE GENOMICS, INC.
 BANDMAN, Olga
 LU, Dyung Aina M.
 YUE, Henry
 TRAN, Bao
 HILLMAN, Jennifer L.
 BAUGHN, Mariah R.
 LAL, Preeti
 TANG, Y. Tom

<120> ISOMERASE PROTEINS

<130> PF-0730 PCT

<140> To Be Assigned

<141> Herewith

<150> 60/149,388

<151> 1999-08-17

<160> 16

<170> PERL Program

<210> 1

<211> 542

<212> PRT

<213> Homo sapiens

<220>

<221> misc_feature

<223> Incyte ID No: 011886CD1

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 Pro Asn Gly Leu Ile Leu Gln Tyr Gly Thr Ala Gly Phe Arg Thr
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 Lys Ala Glu His Leu Asp His Val Met Phe Arg Met Gly Leu Leu
 35 40 45
 Ala Val Leu Arg Ser Lys Gln Thr Lys Ser Thr Ile Gly Val Met
 50 55 60
 Val Thr Ala Ser His Asn Pro Glu Glu Asp Asn Gly Val Lys Leu
 65 70 75
 Val Asp Pro Leu Gly Glu Met Leu Ala Pro Ser Trp Glu Glu His
 80 85 90
 Ala Thr Cys Leu Ala Asn Ala Glu Glu Gln Asp Met Gln Arg Val
 95 100 105
 Leu Ile Asp Ile Ser Glu Lys Glu Ala Val Asn Leu Gln Gln Asp
 110 115 120
 Ala Phe Val Val Ile Gly Arg Asp Thr Arg Pro Ser Ser Glu Lys
 125 130 135
 Leu Ser Gln Ser Val Ile Asp Gly Val Thr Val Leu Gly Gly Gln
 140 145 150
 Phe His Asp Tyr Gly Leu Leu Thr Thr Pro Gln Leu His Tyr Met
 155 160 165
 Val Tyr Cys Arg Asn Thr Gly Gly Arg Tyr Gly Lys Ala Thr Ile
 170 175 180
 Glu Gly Tyr Tyr Gln Lys Leu Ser Lys Ala Phe Val Glu Leu Thr

Lys Gln Ala Ser	185		190		195
Cys Ser Gly Asp Glu Tyr Arg Ser Leu Lys Val	200		205		210
Asp Cys Ala Asn Gly Ile Gly Ala Leu Lys Leu Arg Glu Met Glu	215		220		225
His Tyr Phe Ser Gln Gly Leu Ser Val Gln Leu Phe Asn Asp Gly	230		235		240
Ser Lys Gly Lys Leu Asn His Leu Cys Gly Ala Asp Phe Val Lys	245		250		255
Ser His Gln Lys Pro Pro Gln Gly Met Glu Ile Lys Ser Asn Glu	260		265		270
Arg Cys Cys Ser Phe Asp Gly Asp Ala Asp Arg Ile Val Tyr Tyr	275		280		285
Tyr His Asp Ala Asp Gly His Phe His Leu Ile Asp Gly Asp Lys	290		295		300
Ile Ala Thr Leu Ile Ser Ser Phe Leu Lys Glu Leu Leu Val Glu	305		310		315
Ile Gly Glu Ser Leu Asn Ile Gly Val Val Gln Thr Ala Tyr Ala	320		325		330
Asn Gly Ser Ser Thr Arg Tyr Leu Glu Glu Val Met Lys Val Pro	335		340		345
Val Tyr Cys Thr Lys Thr Gly Val Lys His Leu His His Lys Ala	350		355		360
Gln Glu Phe Asp Ile Gly Val Tyr Phe Glu Ala Asn Gly His Gly	365		370		375
Thr Ala Leu Phe Ser Thr Ala Val Glu Met Lys Ile Lys Gln Ser	380		385		390
Ala Glu Gln Leu Glu Asp Lys Lys Arg Lys Ala Ala Lys Met Leu	395		400		405
Glu Asn Ile Ile Asp Leu Phe Asn Gln Ala Ala Gly Asp Ala Ile	410		415		420
Ser Asp Met Leu Val Ile Glu Ala Ile Leu Ala Leu Lys Gly Leu	425		430		435
Thr Val Gln Gln Trp Asp Ala Leu Tyr Thr Asp Leu Pro Asn Arg	440		445		450
Gln Leu Lys Val Gln Val Ala Asp Arg Arg Val Ile Ser Thr Thr	455		460		465
Asp Ala Glu Arg Gln Ala Val Thr Pro Pro Gly Leu Gln Glu Ala	470		475		480
Ile Asn Asp Leu Val Lys Lys Tyr Lys Leu Ser Arg Ala Phe Val	485		490		495
Arg Pro Ser Gly Thr Glu Asp Val Val Arg Val Tyr Ala Glu Ala	500		505		510
Asp Ser Gln Glu Ser Ala Asp His Leu Ala His Glu Val Ser Leu	515		520		525
Ala Val Phe Gln Leu Ala Gly Gly Ile Gly Glu Arg Pro Gln Pro	530		535		540
Gly Phe					

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<213> Homo sapiens

<220>

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<223> Incyte ID No: 1863189CD1

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Ala Pro Leu Pro Gly	Arg Ala Gly Gly	Ala Ala Ser Gly Gly	Gly Gly
20	25	30	
Gly Asn Ser Trp Asp	Leu Pro Gly Ser His	Val Arg Leu Pro Gly	
35	40	45	
Arg Ala Gln Ser Gly	Thr Arg Gly Gly	Ala Gly Asn Thr Ser	Thr
50	55	60	
Ser Cys Gly Asp Ser	Asn Ser Ile Cys Pro	Ala Pro Ser Thr Met	
65	70	75	
Ser Lys Ala Glu Glu	Ala Lys Lys Leu Ala	Gly Arg Ala Ala Val	
80	85	90	
Glu Asn His Val Arg	Asn Asn Gln Val Leu	Gly Ile Gly Ser Gly	
95	100	105	
Ser Thr Ile Val His	Ala Val Gln Arg Ile	Ala Glu Arg Val Lys	
110	115	120	
Gln Glu Asn Leu Asn	Leu Val Cys Ile Pro	Thr Ser Phe Gln Ala	
125	130	135	
Arg Gln Leu Ile Leu	Gln Tyr Gly Leu Thr	Leu Ser Asp Leu Asp	
140	145	150	
Arg His Pro Glu Ile	Asp Leu Ala Ile Asp	Gly Ala Asp Glu Val	
155	160	165	
Asp Ala Asp Leu Asn	Leu Ile Lys Gly Gly	Gly Gly Cys Leu Thr	
170	175	180	
Gln Glu Lys Ile Val	Ala Gly Tyr Ala Ser	Arg Phe Ile Val Ile	
185	190	195	
Ala Asp Phe Arg Lys	Asp Ser Lys Asn Leu	Gly Asp Gln Trp His	
200	205	210	
Lys Gly Ile Pro Ile	Glu Val Ile Pro Met	Ala Tyr Val Pro Val	
215	220	225	
Ser Arg Ala Val Ser	Gln Lys Phe Gly Gly	Val Val Glu Leu Arg	
230	235	240	
Met Ala Val Asn Lys	Ala Gly Pro Val Val	Thr Asp Asn Gly Asn	
245	250	255	
Phe Ile Leu Asp Trp	Lys Phe Asp Arg Val	His Lys Trp Ser Glu	
260	265	270	
Val Asn Thr Ala Ile	Lys Met Ile Pro Gly	Val Val Asp Thr Gly	
275	280	285	
Leu Phe Ile Asn Met	Ala Glu Arg Val Tyr	Phe Gly Met Gln Asp	
290	295	300	
Gly Ser Val Asn Met	Arg Glu Lys Pro Phe	Cys	
305	310		

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<211> 273

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<213> Homo sapiens

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<223> Incyte ID No: 2088868CD1

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Thr Cys Glu Leu Ala	Glu Val Ala Ala	Glu Val Glu Lys Ser
20	25	30
Ser Asp Gly Pro Gly	Ala Ala Gln Glu Pro	Thr Trp Leu Thr Asp
35	40	45
Val Pro Ala Ala Met	Glu Phe Ile Ala Ala	Thr Glu Val Ala Val
50	55	60

Ile	Gly	Phe	Phe	Gln	Asp	Leu	Glu	Ile	Pro	Ala	Val	Pro	Ile	Leu	
				65					70					75	
His	Ser	Met	Val	Gln	Lys	Phe	Pro	Gly	Val	Ser	Phe	Gly	Ile	Ser	
				80					85					90	
Thr	Asp	Ser	Glu	Val	Leu	Thr	His	Tyr	Asn	Ile	Thr	Gly	Asn	Thr	
				95					100					105	
Ile	Cys	Leu	Phe	Arg	Leu	Val	Asp	Asn	Glu	Gln	Leu	Asn	Leu	Glu	
				110					115					120	
Asp	Glu	Asp	Ile	Glu	Ser	Ile	Asp	Ala	Thr	Lys	Leu	Ser	Arg	Phe	
				125					130					135	
Ile	Glu	Ile	Asn	Ser	Leu	His	Met	Val	Thr	Glu	Tyr	Asn	Pro	Val	
				140					145					150	
Thr	Val	Ile	Gly	Leu	Phe	Asn	Ser	Val	Ile	Gln	Ile	His	Leu	Leu	
				155					160					165	
Leu	Ile	Met	Asn	Lys	Ala	Ser	Pro	Glu	Tyr	Glu	Glu	Asn	Met	His	
				170					175					180	
Arg	Tyr	Gln	Lys	Ala	Ala	Lys	Leu	Phe	Gln	Gly	Lys	Ile	Leu	Phe	
				185					190					195	
Ile	Leu	Val	Asp	Ser	Gly	Met	Lys	Glu	Asn	Gly	Lys	Val	Ile	Ser	
				200					205					210	
Phe	Phe	Lys	Leu	Lys	Glu	Ser	Gln	Leu	Pro	Ala	Leu	Ala	Ile	Tyr	
				215					220					225	
Gln	Thr	Leu	Asp	Asp	Glu	Trp	Asp	Thr	Leu	Pro	Thr	Ala	Glu	Val	
				230					235					240	
Ser	Val	Glu	His	Val	Gln	Asn	Phe	Cys	Asp	Gly	Phe	Leu	Ser	Gly	
				245					250					255	
Lys	Leu	Leu	Lys	Glu	Asn	Arg	Glu	Ser	Glu	Gly	Lys	Thr	Pro	Lys	
				260					265					270	
Val	Glu	Leu													

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 <213> Homo sapiens

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Leu	Ala	Asn	Leu	Gly	Ala	Glu	Cys	Leu	Arg	Met	Leu	Asp	Ser	Gly	
				20					25					30	
Ala	Asp	Tyr	Leu	His	Leu	Asp	Val	Met	Asp	Gly	His	Phe	Val	Pro	
				35					40					45	
Asn	Ile	Thr	Phe	Gly	His	Pro	Val	Val	Glu	Ser	Leu	Arg	Lys	Gln	
				50					55					60	
Leu	Gly	Gln	Asp	Pro	Phe	Phe	Asp	Met	His	Met	Met	Val	Ser	Lys	
				65					70					75	
Pro	Glu	Gln	Trp	Val	Lys	Pro	Met	Ala	Val	Ala	Gly	Ala	Asn	Gln	
				80					85					90	
Tyr	Thr	Phe	His	Leu	Glu	Ala	Thr	Glu	Asn	Pro	Gly	Ala	Leu	Ile	
				95					100					105	
Lys	Asp	Ile	Arg	Glu	Asn	Gly	Met	Lys	Val	Gly	Leu	Ala	Ile	Lys	
				110					115					120	
Pro	Gly	Thr	Ser	Val	Glu	Tyr	Leu	Ala	Pro	Trp	Ala	Asn	Gln	Ile	
				125					130					135	
Asp	Met	Ala	Leu	Val	Met	Thr	Val	Glu	Pro	Gly	Phe	Gly	Gly	Gln	
				140					145					150	

Lys	Phe	Met	Glu	Asp	Met	Met	Pro	Lys	Val	His	Trp	Leu	Arg	Thr	
				155					160					165	
Gln	Phe	Pro	Ser	Leu	Asp	Ile	Glu	Val	Asp	Gly	Gly	Val	Gly	Pro	
				170					175					180	
Asp	Thr	Val	His	Lys	Cys	Ala	Glu	Ala	Gly	Ala	Asn	Met	Ile	Val	
				185					190					195	
Ser	Gly	Ser	Ala	Ile	Met	Arg	Ser	Glu	Asp	Pro	Arg	Ser	Val	Ile	
				200					205					210	
Asn	Leu	Leu	Arg	Asn	Val	Cys	Ser	Glu	Ala	Ala	Gln	Lys	Arg	Ser	
				215					220					225	
Leu	Asp	Arg													

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<213> Homo sapiens

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Met	Gly	Val	Trp	Leu	Asn	Lys	Asp	Asp	Asp	Ile	Arg	Asp	Leu	Lys	
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Arg	Ile	Ile	Leu	Cys	Phe	Leu	Ile	Val	Tyr	Met	Ala	Ile	Leu	Val	
				20					25					30	
Gly	Thr	Asp	Gln	Asp	Phe	Tyr	Ser	Leu	Leu	Gly	Val	Ser	Lys	Thr	
				35					40					45	
Ala	Ser	Ser	Arg	Glu	Ile	Arg	Gln	Ala	Phe	Lys	Lys	Leu	Ala	Leu	
				50					55					60	
Lys	Leu	His	Pro	Asp	Lys	Asn	Pro	Asn	Asn	Pro	Asn	Ala	His	Gly	
				65					70					75	
Asn	Phe	Leu	Lys	Ile	Asn	Arg	Ala	Tyr	Glu	Val	Leu	Lys	Asp	Glu	
				80					85					90	
Asp	Leu	Arg	Lys	Lys	Tyr	Asp	Lys	Tyr	Gly	Glu	Lys	Gly	Leu	Glu	
				95					100					105	
Asp	Asn	Gln	Gly	Gly	Gln	Tyr	Glu	Ser	Trp	Asn	Tyr	Tyr	Arg	Tyr	
				110					115					120	
Asp	Phe	Gly	Ile	Tyr	Asp	Asp	Asp	Pro	Glu	Ile	Ile	Thr	Leu	Glu	
				125					130					135	
Arg	Arg	Glu	Phe	Asp	Ala	Ala	Val	Asn	Ser	Gly	Glu	Leu	Trp	Phe	
				140					145					150	
Val	Asn	Phe	Tyr	Ser	Pro	Gly	Cys	Ser	His	Cys	His	Asp	Leu	Ala	
				155					160					165	
Pro	Thr	Trp	Arg	Asp	Phe	Ala	Lys	Glu	Val	Asp	Gly	Leu	Leu	Arg	
				170					175					180	
Ile	Gly	Ala	Val	Asn	Cys	Gly	Asp	Asp	Arg	Met	Leu	Cys	Arg	Met	
				185					190					195	
Lys	Gly	Val	Asn	Ser	Tyr	Pro	Ser	Leu	Phe	Ile	Phe	Arg	Ser	Gly	
				200					205					210	
Met	Ala	Pro	Val	Lys	Tyr	His	Gly	Asp	Arg	Ser	Lys	Glu	Ser	Leu	
				215					220					225	
Val	Ser	Phe	Ala	Met	Gln	His	Val	Arg	Ser	Thr	Val	Thr	Glu	Leu	
				230					235					240	
Trp	Thr	Gly	Asn	Phe	Val	Asn	Ser	Ile	Gln	Thr	Ala	Phe	Ala	Ala	
				245					250					255	
Gly	Ile	Gly	Trp	Leu	Ile	Thr	Phe	Cys	Ser	Lys	Gly	Gly	Asp	Cys	
				260					265					270	
Leu	Thr	Ser	Gln	Thr	Arg	Leu	Arg	Leu	Ser	Gly	Met	Leu	Asp	Gly	
				275					280					285	

Leu	Val	Asn	Val	Gly	Trp	Met	Asp	Cys	Ala	Thr	Gln	Asp	Asn	Leu
				290					295					300
Cys	Lys	Ser	Leu	Asp	Ile	Thr	Thr	Ser	Thr	Thr	Ala	Tyr	Phe	Pro
				305					310					315
Pro	Gly	Ala	Thr	Leu	Asn	Asn	Lys	Glu	Lys	Asn	Ser	Ile	Leu	Phe
				320					325					330
Leu	Asn	Ser	Leu	Asp	Ala	Lys	Glu	Ile	Tyr	Leu	Glu	Val	Ile	His
				335					340					345
Asn	Leu	Pro	Asp	Phe	Glu	Leu	Leu	Ser	Ala	Asn	Thr	Leu	Glu	Asp
				350					355					360
Arg	Leu	Ala	His	His	Arg	Trp	Leu	Leu	Phe	Phe	His	Phe	Gly	Lys
				365					370					375
Asn	Glu	Asn	Ser	Asn	Asp	Pro	Glu	Leu	Lys	Lys	Leu	Lys	Thr	Leu
				380					385					390
Leu	Lys	Asn	Asp	His	Ile	Gln	Val	Gly	Arg	Phe	Asp	Cys	Ser	Ser
				395					400					405
Ala	Pro	Asp	Ile	Cys	Ser	Asn	Leu	Tyr	Val	Phe	Gln	Pro	Ser	Leu
				410					415					420
Ala	Val	Phe	Lys	Gly	Gln	Gly	Thr	Lys	Glu	Tyr	Glu	Ile	His	His
				425					430					435
Gly	Lys	Lys	Ile	Leu	Tyr	Asp	Ile	Leu	Ala	Phe	Ala	Lys	Glu	Ser
				440					445					450
Val	Asn	Ser	His	Val	Thr	Thr	Leu	Gly	Pro	Gln	Asn	Phe	Pro	Ala
				455					460					465
Asn	Asp	Lys	Glu	Pro	Trp	Leu	Val	Asp	Phe	Phe	Ala	Pro	Trp	Cys
				470					475					480
Pro	Pro	Cys	Arg	Ala	Leu	Leu	Pro	Glu	Leu	Arg	Arg	Ala	Ser	Asn
				485					490					495
Leu	Leu	Tyr	Gly	Gln	Leu	Lys	Phe	Gly	Thr	Leu	Asp	Cys	Thr	Val
				500					505					510
His	Glu	Gly	Leu	Cys	Asn	Met	Tyr	Asn	Ile	Gln	Ala	Tyr	Pro	Thr
				515					520					525
Thr	Val	Val	Phe	Asn	Gln	Ser	Asn	Ile	His	Glu	Tyr	Glu	Gly	His
				530					535					540
His	Ser	Ala	Glu	Gln	Ile	Leu	Glu	Phe	Ile	Glu	Asp	Leu	Met	Asn
				545					550					555
Pro	Ser	Val	Val	Ser	Leu	Thr	Pro	Thr	Thr	Phe	Asn	Glu	Leu	Val
				560					565					570
Thr	Gln	Arg	Lys	His	Asn	Glu	Val	Trp	Met	Val	Asp	Phe	Tyr	Ser
				575					580					585
Pro	Trp	Cys	His	Pro	Cys	Gln	Val	Leu	Met	Pro	Glu	Trp	Lys	Arg
				590					595					600
Met	Ala	Arg	Thr	Leu	Thr	Gly	Leu	Ile	Asn	Val	Gly	Ser	Ile	Asp
				605					610					615
Cys	Gln	Gln	Tyr	His	Ser	Phe	Cys	Ala	Gln	Glu	Asn	Val	Gln	Arg
				620					625					630
Tyr	Pro	Glu	Ile	Arg	Phe	Phe	Pro	Pro	Lys	Ser	Asn	Lys	Ala	Tyr
				635					640					645
Gln	Tyr	His	Ser	Tyr	Asn	Gly	Trp	Asn	Arg	Asp	Ala	Tyr	Ser	Leu
				650					655					660
Arg	Ile	Trp	Gly	Leu	Gly	Phe	Leu	Pro	Gln	Val	Ser	Thr	Asp	Leu
				665					670					675
Thr	Pro	Gln	Thr	Phe	Ser	Glu	Lys	Val	Leu	Gln	Gly	Lys	Asn	His
				680					685					690
Trp	Val	Ile	Asp	Phe	Tyr	Ala	Pro	Trp	Cys	Gly	Pro	Cys	Gln	Asn
				695					700					705
Phe	Ala	Pro	Glu	Phe	Glu	Leu	Leu	Ala	Arg	Met	Ile	Lys	Gly	Lys
				710					715					720
Val	Lys	Ala	Gly	Lys	Val	Asp	Cys	Gln	Ala	Tyr	Ala	Gln	Thr	Cys
				725					730					735
Gln	Lys	Ala	Gly	Ile	Arg	Ala	Tyr	Pro	Thr	Val	Lys	Phe	Tyr	Phe
				740					745					750
Tyr	Glu	Arg	Ala	Lys	Arg	Asn	Phe	Gln	Glu	Glu	Gln	Ile	Asn	Thr

	755		760		765
Arg Asp Ala Lys	Ala Ile Ala Ala Leu	Ile Ser Glu Lys Leu	Glu		
	770		775		780
Thr Leu Arg Asn	Gln Gly Lys Arg Asn	Lys Asp Glu Leu			
	785		790		

<210> 6
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<400> 6

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Leu	Tyr	Thr	Glu	Glu	Arg	Pro	Arg	Ala	Cys	Leu	Asn	Phe	Leu	Lys
				20					25					30
Leu	Cys	Lys	Ile	Lys	Tyr	Tyr	Asn	Tyr	Cys	Leu	Ile	His	Asn	Val
				35					40					45
Gln	Arg	Asp	Phe	Ile	Ile	Gln	Thr	Gly	Asp	Pro	Thr	Gly	Thr	Gly
				50					55					60
Arg	Gly	Gly	Glu	Ser	Ile	Phe	Gly	Gln	Leu	Tyr	Gly	Asp	Gln	Ala
				65					70					75
Ser	Phe	Phe	Glu	Ala	Glu	Lys	Val	Pro	Arg	Ile	Lys	His	Lys	Lys
				80					85					90
Lys	Gly	Thr	Val	Ser	Met	Val	Asn	Asn	Gly	Ser	Asp	Gln	His	Gly
				95					100					105
Ser	Gln	Phe	Leu	Ile	Thr	Thr	Gly	Glu	Asn	Leu	Asp	Tyr	Leu	Asp
				110					115					120
Gly	Val	His	Thr	Val	Phe	Gly	Glu	Val	Thr	Glu	Gly	Met	Asp	Ile
				125					130					135
Ile	Lys	Lys	Ile	Asn	Glu	Thr	Phe	Val	Asp	Lys	Asp	Phe	Val	Pro
				140					145					150
Tyr	Gln	Asp	Ile	Arg	Ile	Asn	His	Thr	Val	Ile	Leu	Asp	Asp	Pro
				155					160					165
Phe	Asp	Asp	Pro	Pro	Asp	Leu	Leu	Ile	Pro	Asp	Arg	Ser	Pro	Glu
				170					175					180
Pro	Thr	Arg	Glu	Gln	Leu	Asp	Ser	Gly	Arg	Ile	Gly	Ala	Asp	Glu
				185					190					195
Glu	Ile	Asp	Asp	Phe	Lys	Gly	Arg	Ser	Ala	Glu	Glu	Val	Glu	Glu
				200					205					210
Ile	Lys	Ala	Glu	Lys	Glu	Ala	Lys	Thr	Gln	Ala	Ile	Leu	Leu	Glu
				215					220					225
Met	Val	Gly	Asp	Leu	Pro	Asp	Ala	Asp	Ile	Lys	Pro	Pro	Glu	Asn
				230					235					240
Val	Leu	Phe	Val	Cys	Lys	Leu	Asn	Pro	Val	Thr	Thr	Asp	Glu	Asp
				245					250					255
Leu	Glu	Ile	Ile	Phe	Ser	Arg	Phe	Gly	Pro	Ile	Arg	Ser	Cys	Glu
				260					265					270
Val	Ile	Arg	Asp	Trp	Lys	Thr	Gly	Glu	Ser	Leu	Cys	Tyr	Ala	Phe
				275					280					285
Ile	Glu	Phe	Glu	Lys	Glu	Glu	Asp	Cys	Glu	Lys	Ala	Phe	Phe	Lys
				290					295					300
Met	Asp	Asn	Val	Leu	Ile	Asp	Asp	Arg	Arg	Ile	His	Val	Asp	Phe
				305					310					315
Ser	Gln	Ser	Val	Ala	Lys	Val	Lys	Trp	Lys	Gly	Lys	Gly	Gly	Lys
				320					325					330

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Tyr Thr Lys Ser Asp Phe Lys Glu Tyr Glu Lys Glu Gln Asp Lys
335 340 345
Pro Pro Asn Leu Val Leu Lys Asp Lys Val Lys Pro Lys Gln Asp
350 355 360
Thr Lys Tyr Asp Leu Ile Leu Asp Glu Gln Ala Glu Asp Ser Lys
365 370 375
Ser Ser His Ser His Thr Ser Lys Lys His Lys Lys Lys Thr His
380 385 390
His Cys Ser Glu Glu Lys Glu Asp Glu Asp Tyr Met Pro Ile Lys
395 400 405
Asn Thr Asn Gln Asp Ile Tyr Arg Glu Met Gly Phe Gly His Tyr
410 415 420
Glu Glu Glu Glu Ser Cys Trp Glu Lys Gln Lys Ser Glu Lys Arg
425 430 435
Asp Arg Thr Gln Asn Arg Ser Arg Ser Arg Ser Arg Glu Arg Asp
440 445 450
Gly His Tyr Ser Asn Ser His Lys Ser Lys Tyr Gln Thr Asp Leu
455 460 465
Tyr Glu Arg Glu Arg Ser Lys Lys Arg Asp Arg Ser Arg Ser Pro
470 475 480
Lys Lys Ser Lys Asp Lys Glu Lys Ser Lys Tyr Arg
485 490

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<210> 7

<211> 160

<212> PRT

<213> Homo sapiens

<220>

<221> misc_feature

<223> Incyte ID No: 3817050CD1

<400> 7

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Met Val Ile Pro Thr Val Pro Phe Asn Ile Thr Ile Asn Ser Lys
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Pro Leu Gly His Ile Ser Phe Gln Leu Phe Ala Asp Lys Phe Pro
20 25 30
Lys Thr Gly Glu Asn Phe His Thr Leu Asn Asn Lys Asp Lys Gly
35 40 45
Phe Gly Ser Cys Phe His Arg Ile Ile Pro Glu Phe Ile Cys Gln
50 55 60
Gly Asp Asp Phe Thr Pro His Asn Gly Ile Gly Gly Lys Ser Ile
65 70 75
Tyr Gly Asp Lys Phe Asp Asp Lys Asn Phe Ile Val Lys His Thr
80 85 90
Gly Leu Gly Ile Leu Ser Met Ala Asn Ala Ala Pro Lys Thr Asn
95 100 105
Glu Ser Gln Phe Phe Ile Cys Thr Ala Met Ala Lys Trp Trp Asp
110 115 120
Gly Lys His Val Ile Phe Gly Arg Val Lys Glu Gly Met Asn Ile
125 130 135
Val Glu Ala Met Glu Cys Phe Gly Ser Arg Asn Gly Lys Thr Ser
140 145 150
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 Ser Pro Gln Pro Pro Gly Tyr Leu Thr Phe Phe Thr Ser Ala Leu
 35 40 45
 His Ser Leu Lys Lys Asp Tyr Leu Gly Thr Val Arg Phe Gly Val
 50 55 60
 Ile Thr Asn Lys His Leu Ala Lys Leu Val Ser Leu Val His Ser
 65 70 75
 Gly Ser Val Tyr Leu His Arg His Phe Asn Thr Ser Leu Val Phe
 80 85 90
 Pro Arg Glu Val Leu Asn Tyr Thr Ala Glu Asn Ile Cys Lys Trp
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 Ala Leu Glu Asn Gln Glu Thr Leu Phe Arg Trp Leu Arg Pro His
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 Gly Gly Lys Ser Leu Leu Leu Asn Asn Glu Leu Lys Lys Gly Pro
 125 130 135
 Ala Leu Phe Leu Phe Ile Pro Phe Asn Pro Leu Ala Glu Ser His
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 Pro Leu Ile Asp Glu Ile Thr Glu Val Ala Leu Glu Tyr Asn Asn
 155 160 165
 Cys His Gly Asp Gln Val Val Glu Arg Leu Leu Gln His Leu Arg
 170 175 180
 Arg Val Asp Ala Pro Val Leu Glu Ser Leu Ala Leu Glu Val Pro
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 Ala Gln Leu Pro Asp Pro Pro Thr Ile Thr Ala Ser Pro Cys Cys
 200 205 210
 Asn Thr Val Val Leu Pro Gln Trp His Ser Phe Ser Arg Thr His
 215 220 225
 Asn Val Cys Glu Leu Cys Val Asn Gln Thr Ser Gly Gly Met Lys
 230 235 240
 Pro Ser Ser Val Ser Val Pro Gln Cys Ser Phe Phe Glu Met Ala
 245 250 255
 Ala Ala Leu Asp Ser Phe Tyr Leu Lys Glu Gln Thr Phe Tyr His
 260 265 270
 Val Ala Ser Asp Ser Ile Glu Cys Ser Asn Phe Leu Thr Ser Tyr
 275 280 285
 Ser Pro Phe Ser Tyr Tyr Thr Ala Cys Cys Arg Thr Ile Ser Arg
 290 295 300
 Gly Val Ser Gly Phe Ile Asp Ser Glu Gln Gly Val Phe Glu Ala
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 Pro Thr Val Ala Phe Ser Ser Leu Glu Lys Lys Cys Glu Val Asp
 320 325 330
 Ala Pro Ser Ser Val Pro His Ile Glu Glu Asn Arg Tyr Leu Phe
 335 340 345
 Pro Glu Val Asp Met Thr Ser Thr Asn Phe Thr Gly Leu Ser Cys
 350 355 360
 Arg Thr Asn Lys Thr Leu Asn Ile Tyr Leu Leu Asp Ser Asn Leu
 365 370 375
 Phe Trp Leu Tyr Ala Glu Arg Leu Gly Ala Pro Ser Ser Thr Gln
 380 385 390
 Val Lys Glu Phe Ala Ala Ile Val Asp Val Lys Glu Glu Ser His
 395 400 405

Tyr	Ile	Leu	Asp	Pro	Lys	Gln	Ala	Leu	Met	Lys	Leu	Thr	Leu	Glu	
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Ser	Phe	Ile	Gln	Asn	Phe	Ser	Val	Leu	Tyr	Ser	Pro	Leu	Lys	Arg	
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His	Leu	Ile	Gly	Ser	Gly	Ser	Ala	Gln	Phe	Pro	Ser	Gln	His	Leu	
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Ile	Thr	Glu	Val	Thr	Thr	Asp	Thr	Phe	Trp	Glu	Val	Val	Leu	Gln	
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Lys	Gln	Asp	Val	Leu	Leu	Leu	Tyr	Tyr	Ala	Pro	Trp	Cys	Gly	Phe	
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Pro	Met	Asp	Thr	Phe	Thr	Val	Ala	Arg	Ile	Asp	Val	Ser	Gln	Asn	
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Phe	Phe	Pro	Cys	Asn	Arg	Lys	Asp	Leu	Ser	Val	Lys	Tyr	Pro	Glu	
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Ile	Ser	His	Leu	Glu	Arg	Glu	Ile	Gln	Lys	Leu	Arg	Ala	Glu	Ile	
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Leu	Glu	Glu	Gln	His	Ser	Leu	Leu	His	Ala	His	Ser	Glu	Gln	Leu	
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Gln	Ala	Leu	Tyr	Glu	Gln	Lys	Thr	Arg	Glu	Leu	Gln	Glu	Leu	Ala	
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Arg	Lys	Leu	Gln	Glu	Leu	Ala	Asp	Ala	Ser	Glu	Asn	Leu	Leu	Thr	
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Glu	Asn	Thr	Trp	Leu	Lys	Ile	Leu	Val	Ala	Thr	Met	Glu	Arg	Lys	
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Leu	Glu	Gly	Arg	Asp	Gly	Ala	Glu	Ser	Leu	Ala	Ala	Gln	Arg	Glu	
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Val	His	Pro	Lys	Gln	Pro	Glu	Pro	Ser	Ala	Thr	Pro	Gln	Leu	Pro	
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Gly	Ser	Ser	Pro	Pro	Pro	Ala	Asn	Val	Ser	Ala	Thr	Leu	Val	Ser	
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 <213> Homo sapiens

<220>
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gtgaa	atggt	ggcaccatcc	tgggaggaac	atgccacctg	tttagcaa	at	360
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taata	gatgg	tgtgactgtt	ctaggagggtc	aattccatga	ttatggcttg	ttaacaacac	540
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<211> 1823

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

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<213> Homo sapiens

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<213> Homo sapiens

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<213> Homo sapiens

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